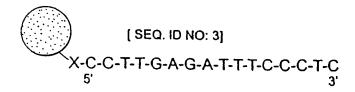
PCT/US01/25237

1/64

FIG. 1



G-G-A-A-C-T-C-T-A-A-A-G-G-G-A-G-X
[SEQ. ID NO: 4]



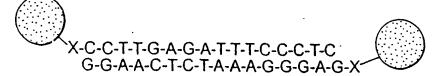


FIG. 2

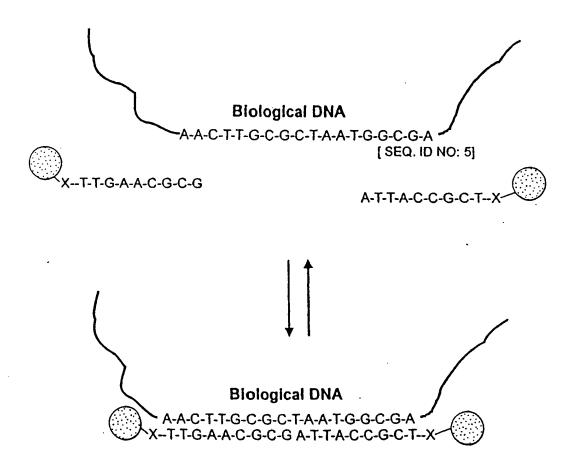
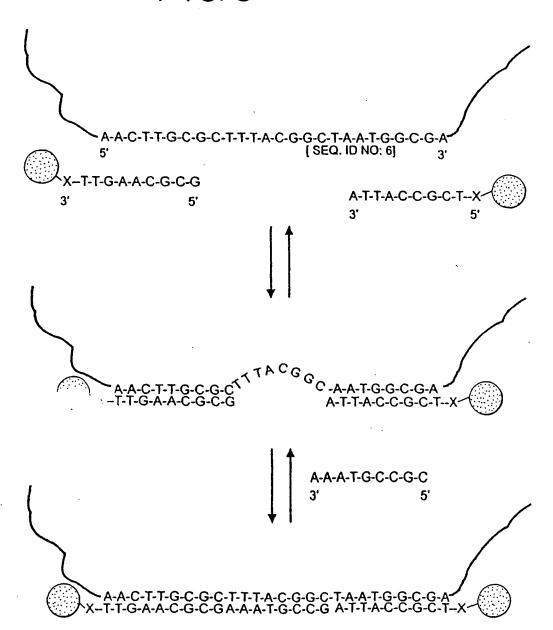
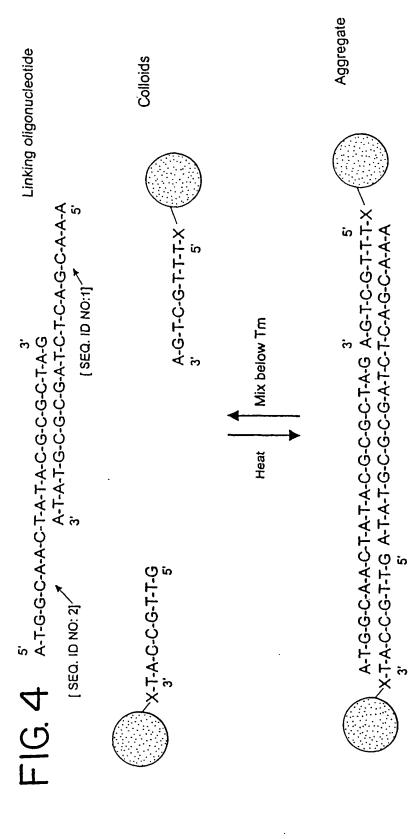


FIG. 3





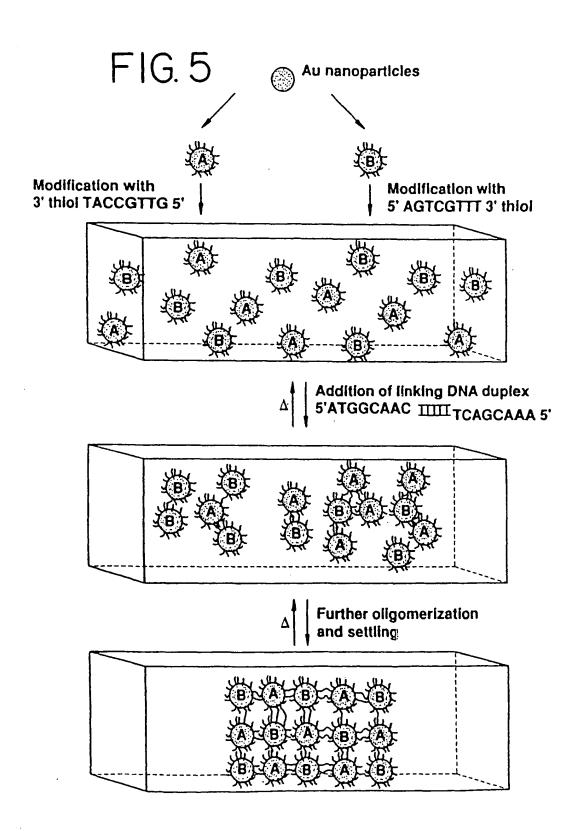
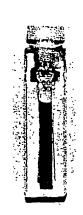
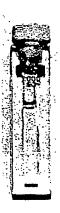
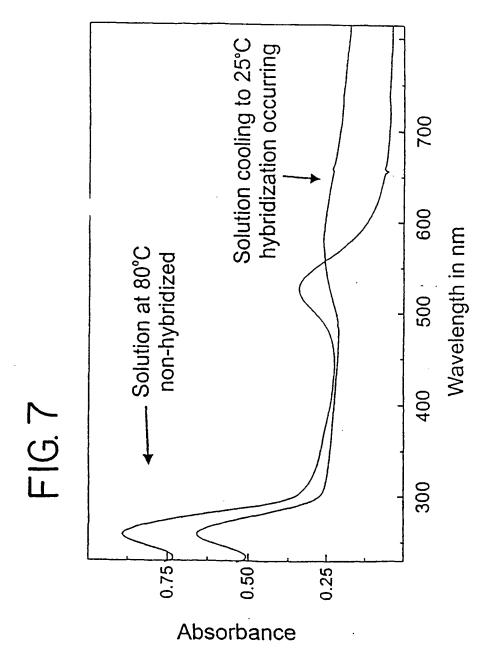


FIG.6A FIG.6B FIG.6C









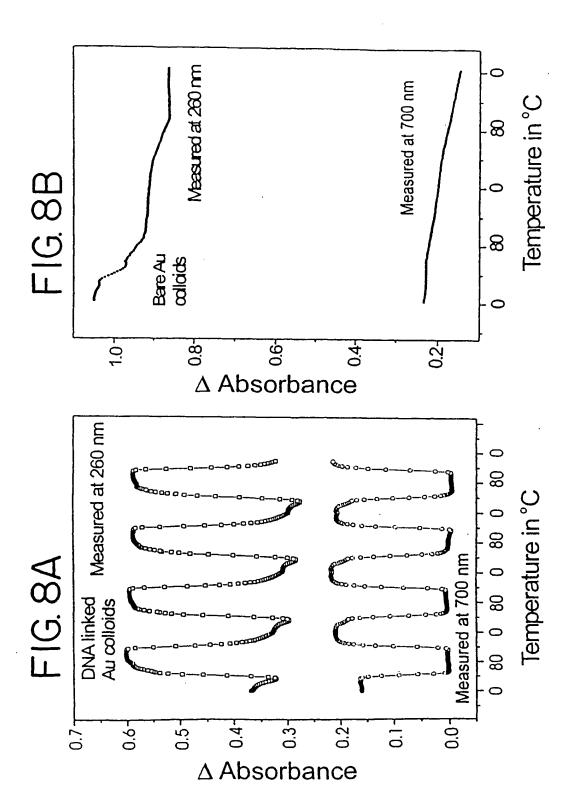


FIG. 9A

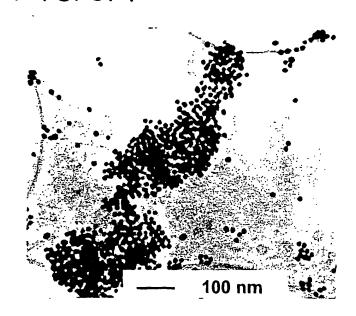
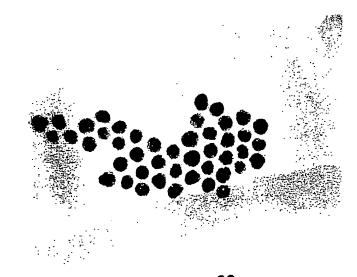
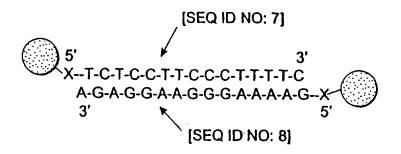
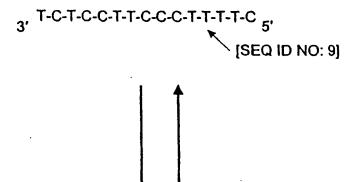


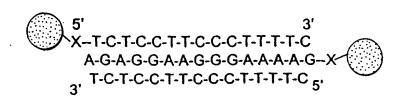
FIG.9B



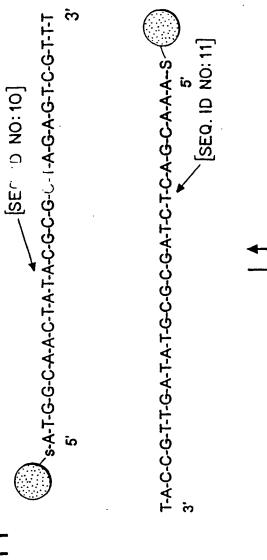
· 33 nm







F1G. |



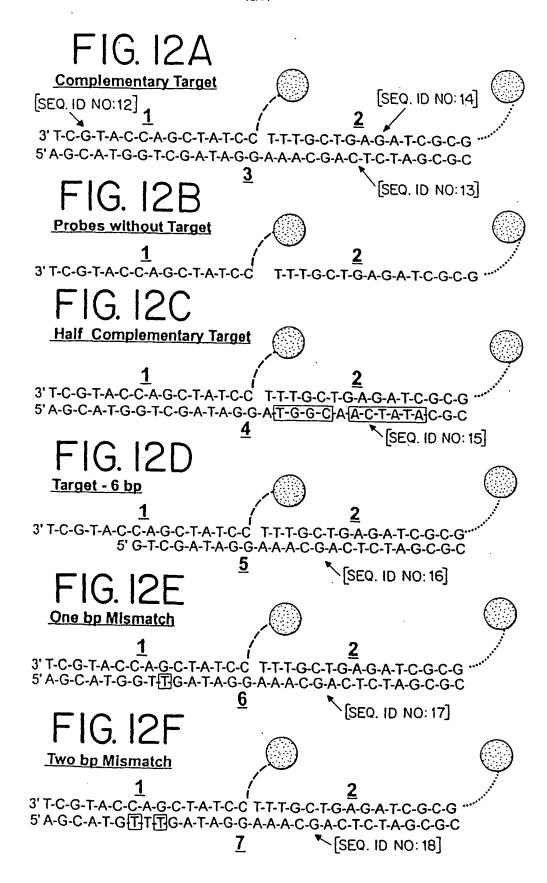


FIG. 13A

transparent substrate

Modified DNA chemisorbed onto solld substrate

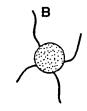
YA,

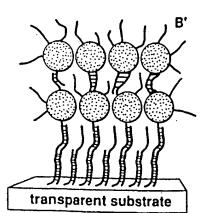
Analyte DNA

transparent substrate

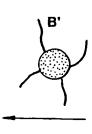
Analyte DNA hybridized onto substrate

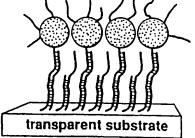
DNA modified colloids



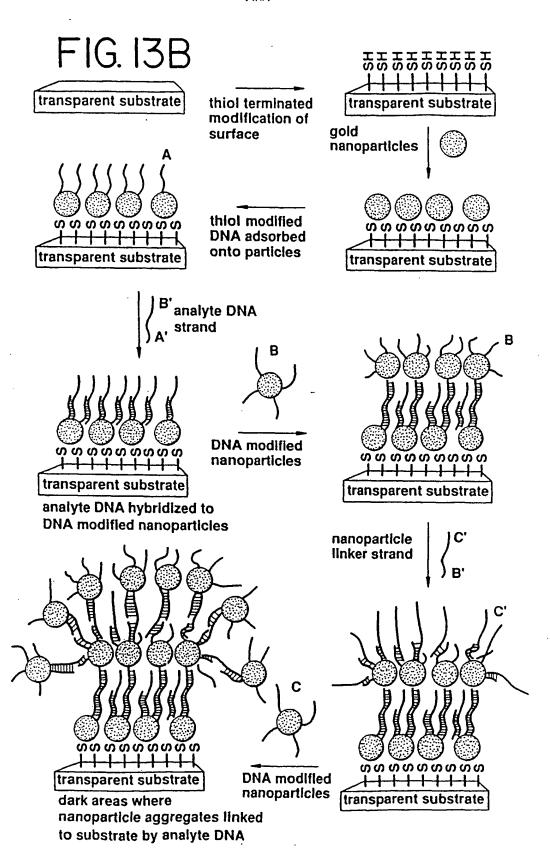


Dark areas where nanoparticle aggregates are linked to substrate surface by analyte DNA





DNA modified colloids hybridized to bound analyte DNA



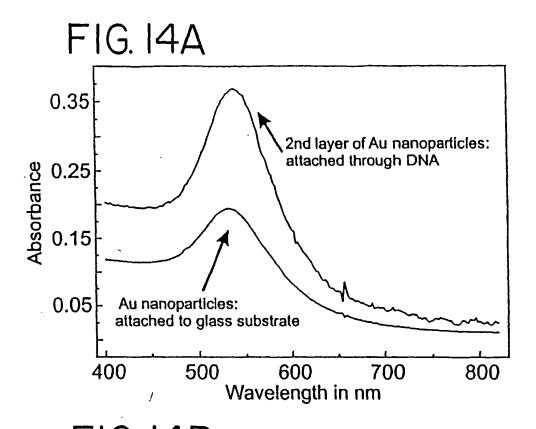


FIG. 14B

0.34

First Derivative: Tm=62 C

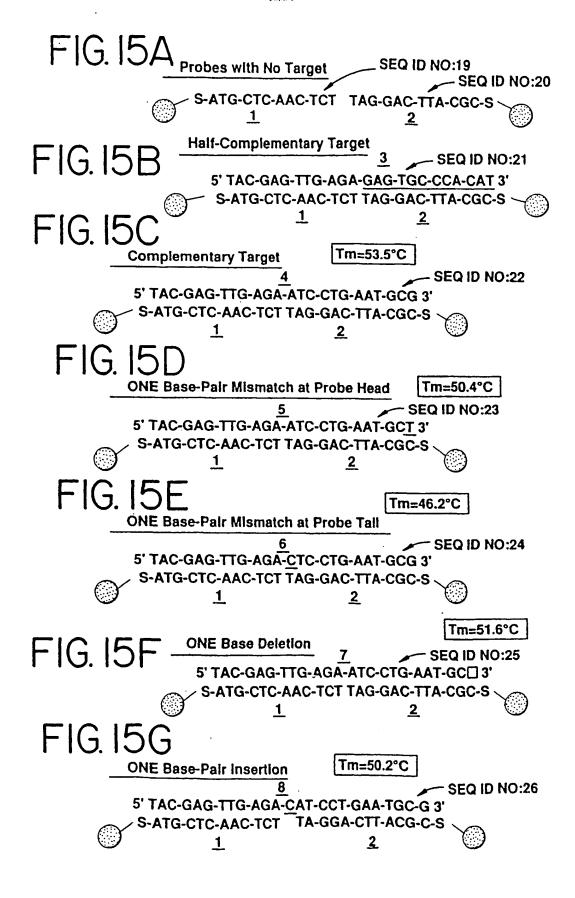
0.26

0.22

0.20

40

Temperature in °C



-16.16A

24 Base Template

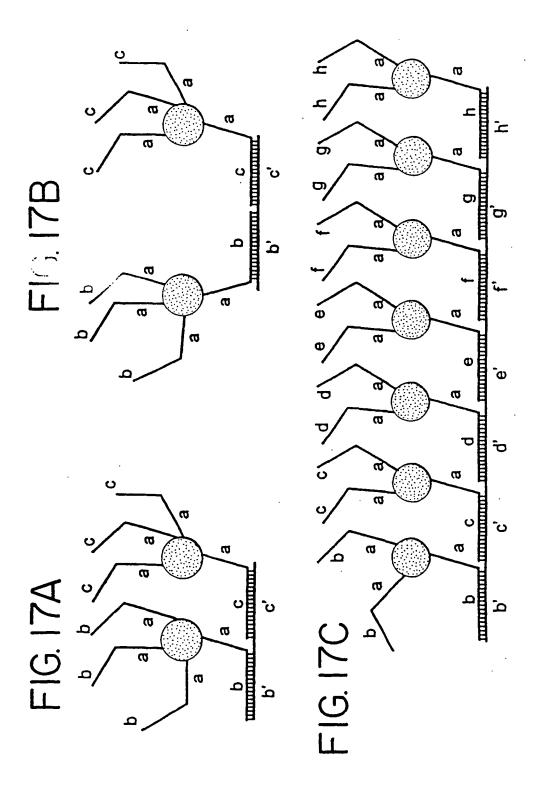
FIG. 16B

48 Base Template with Complementary 24 Base Filler

5' TAC-GAG-TTG-AGA-CCG-TTA-AGA-CGA-GGC-AAT-CAT-GCA-ATC-CTG-AAT-GCG 3' > S-ATG-CTC-AAC-TCT GGC-AAT-TCT-GCT-CCG-TTA-GTA-CGT TAG-GAC-TTA-CGC-S

FIG. 16(

72 Base Template with Complementary 48 Base Filler



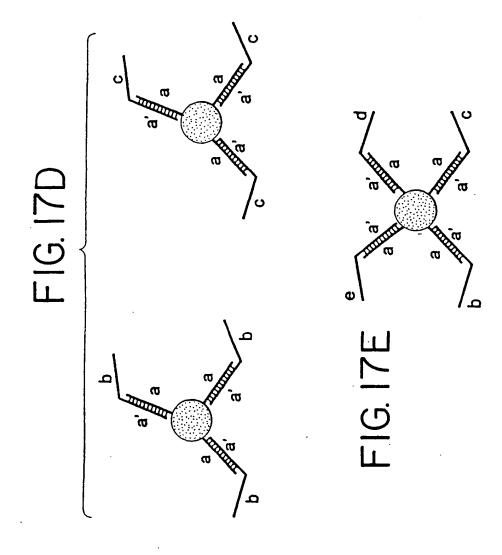
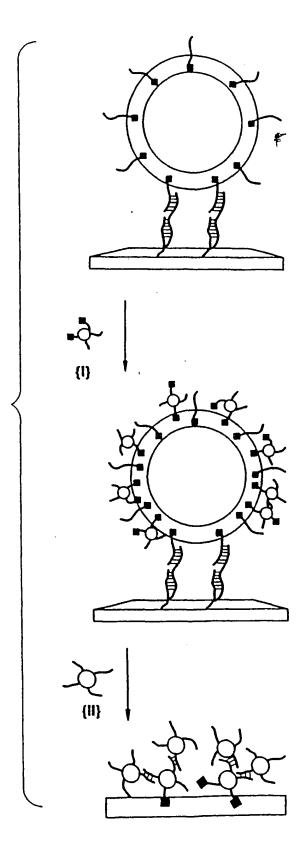


FIG. 18



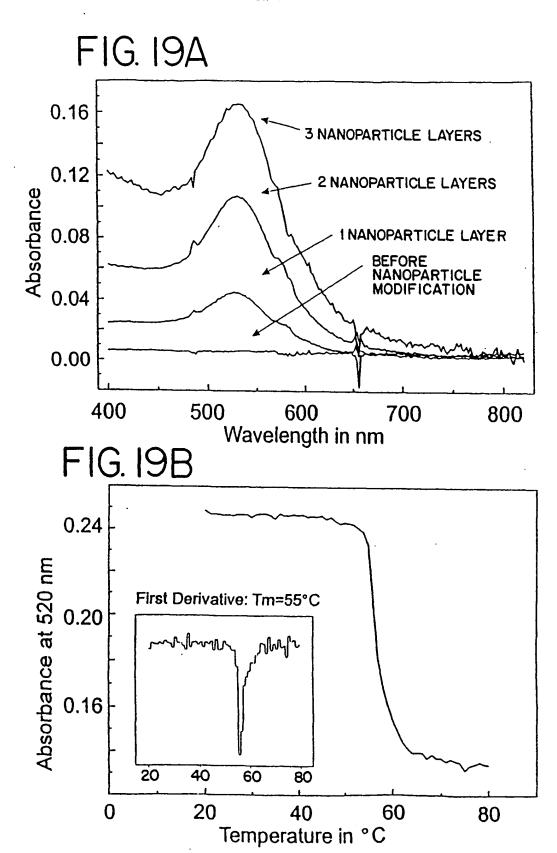
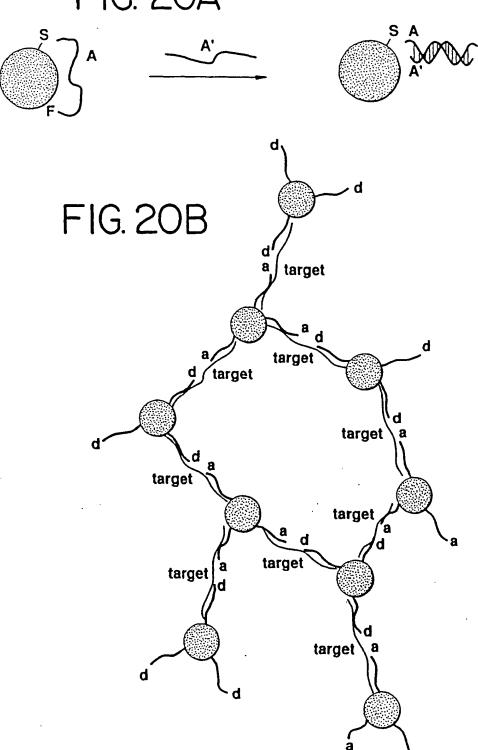
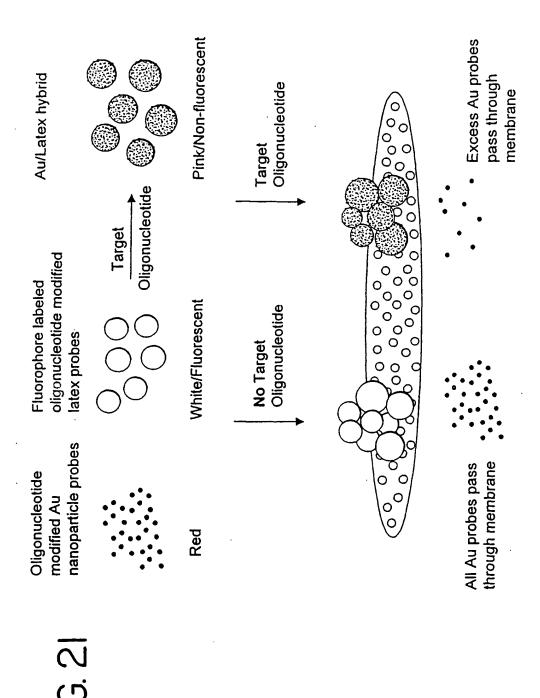
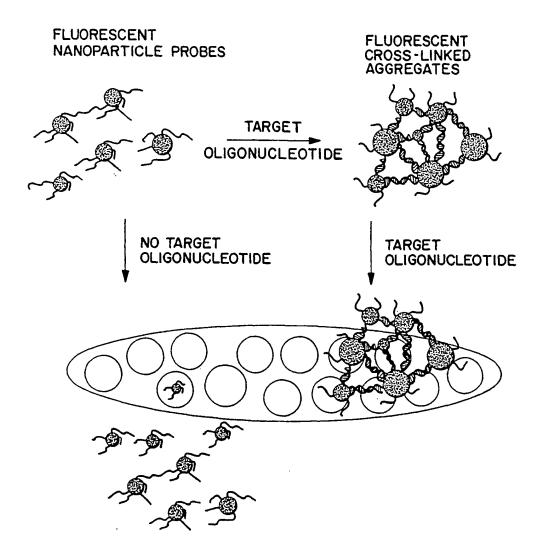


FIG. 20A







THE FLWORESCENT NANOPARTICLE PROBES PASS THROUGH THE MEMBRANE

THE FLUORESCENT CROSS-LINKED AGGREGATES ARE RETAINED BY THE MEMBRANE

Anthrax PCR Product

5'G GCG GAT GAG TCA GTA GTT AAG GAG GCT CAT AGA GAA GTA ATT AAT 3'C CGC CTA CTC AGT CAT CAA TTC CTC CGA GTA TCT CTT CAT TAA TTA

TCG TCA ACA <u>GAG GGA TTA TTG TTA AAT ATT GAT AAG GAT</u> ATA AGA AAA AGC AGT TGT CTC CCT AAT AAC AAT TTA TAA CTA TTC CTA TAT TCT TTT

ATA TTA TCC AGG GTT ATA TTG TAG AAA TTG AAG ATA CTG AAG GGC TT 3' TAT AAT AGG TCC CAA TAT AAC ATC TTT AAC TTC TAT GAC TTC CCG AA 5'

141 mer Anthrax PCR product [SEQ ID NO:36]

3' CTC CCT AAT AAC AAT

[SEQ ID NO:37]

3' TTA TAA CTA TTC CTA -

[SEQ ID NO:38]

Oligonucleotide-Nanoparticle Probes

Blocker Oligonucleotides

3' C CGC CTA CTC AGT CAT CAA TTC CTC CGA GT

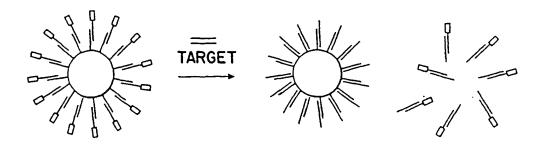
3' A TCT CTT CAT TAA TTA AGC AGT TGT

3' TAT TCT TTT TAT AAT AGG TCC CAA TAT

3' AAC ATC TTT AAC TTC TAT GAC TTC CCG AA

[SEQ ID NO:40]

SEQ ID NO:41]



SATELLITE PROBE

DETECTION SIGNAL

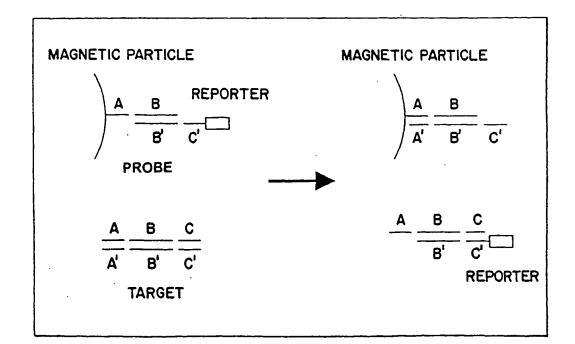
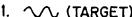
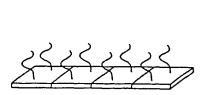


FIG. 25A (TARGET)







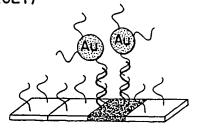
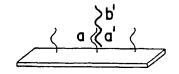
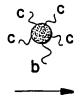


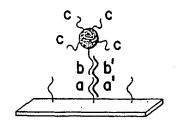
FIG. 25B

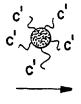


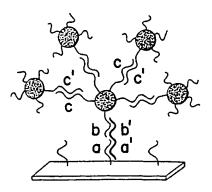














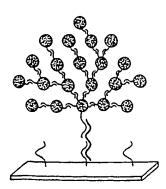


FIG. 26A

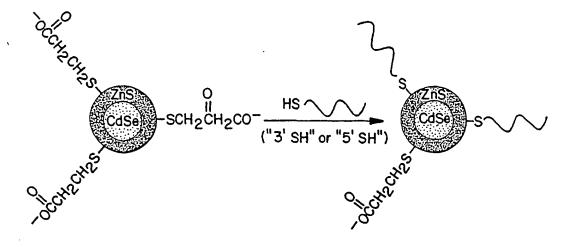
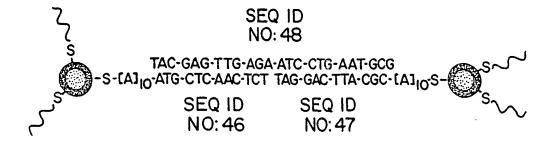


FIG. 26B



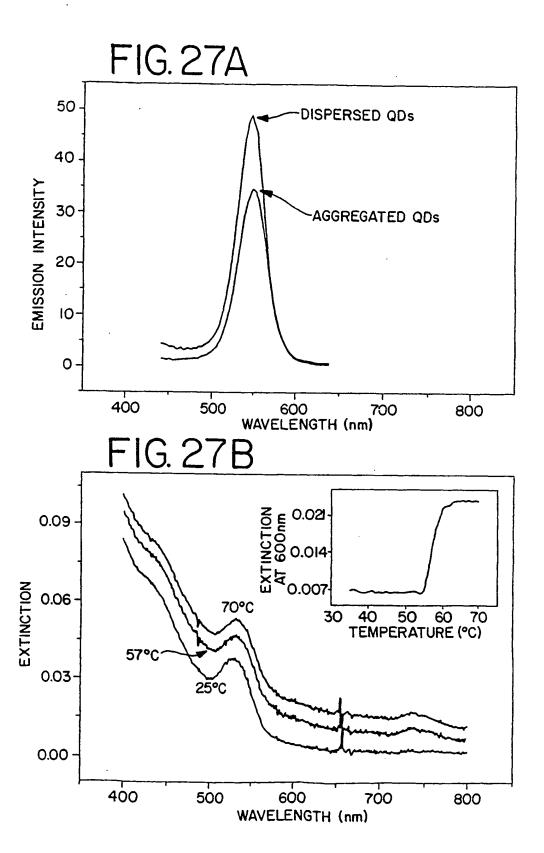


FIG. 27C

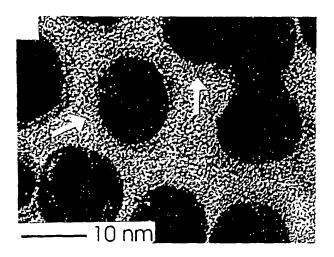


FIG. 27D

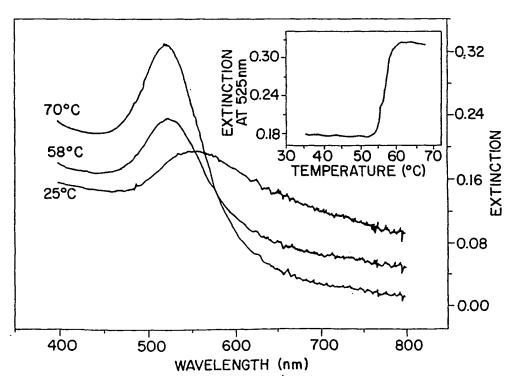


FIG. 28A

FIG. 28B

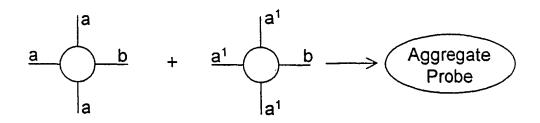


FIG. 28C

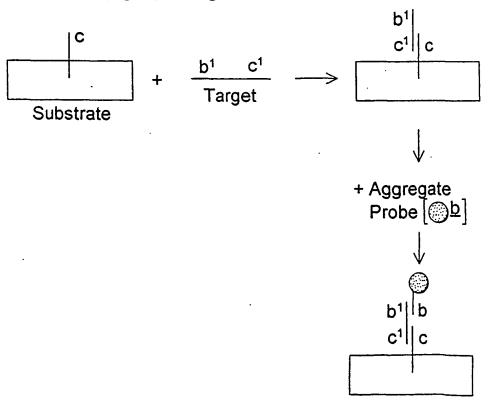


FIG. 28D

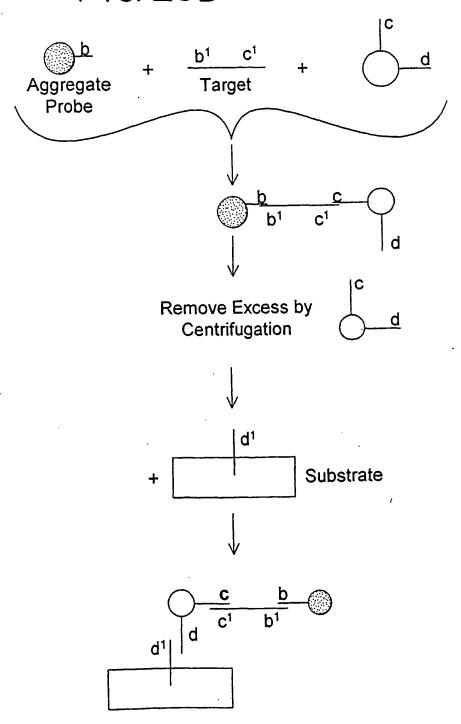
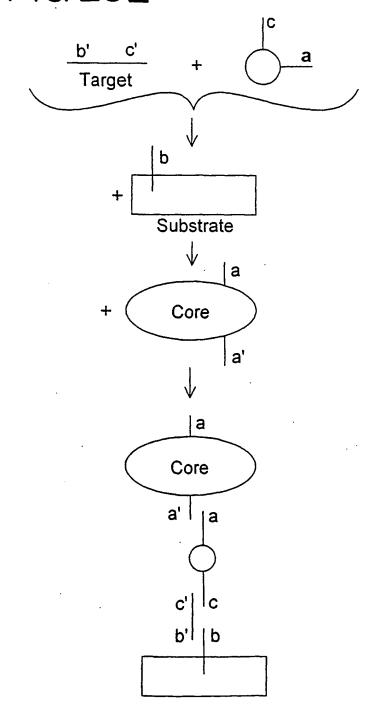
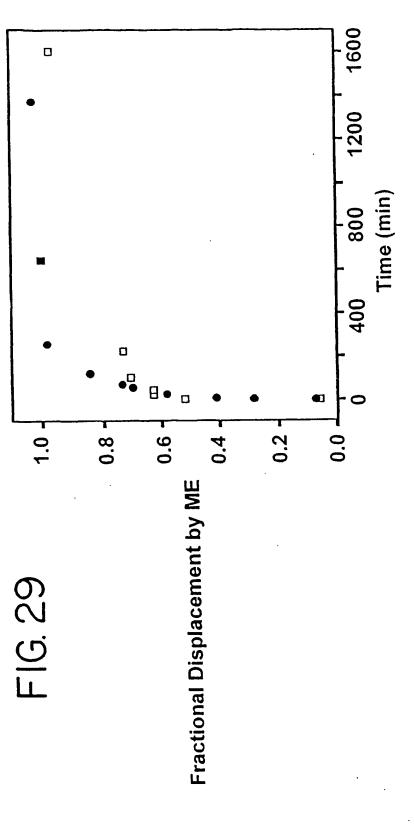
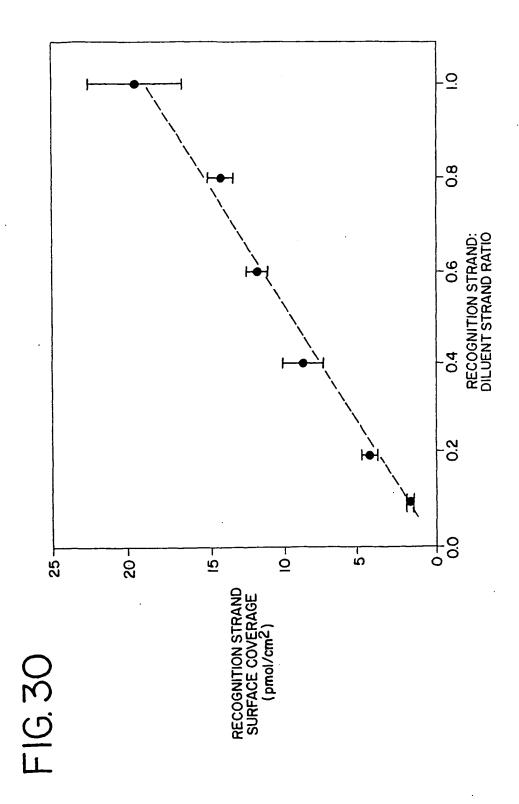
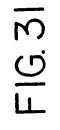


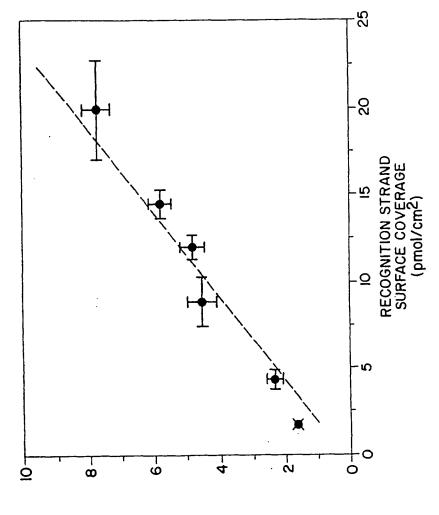
FIG. 28E











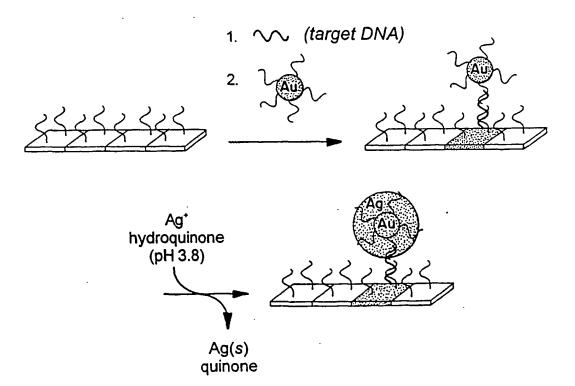
[SEQIDNO:56]

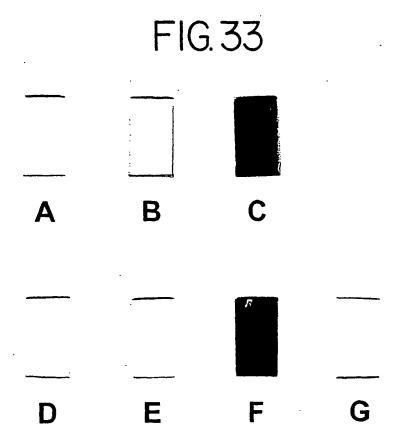
5' GGA T**T**A TTG TTA--AAT ATT GAT AAG GAT 3'

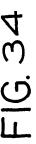
CCT A**N**T AAC AAT TTA TAA CTA TTC CTA

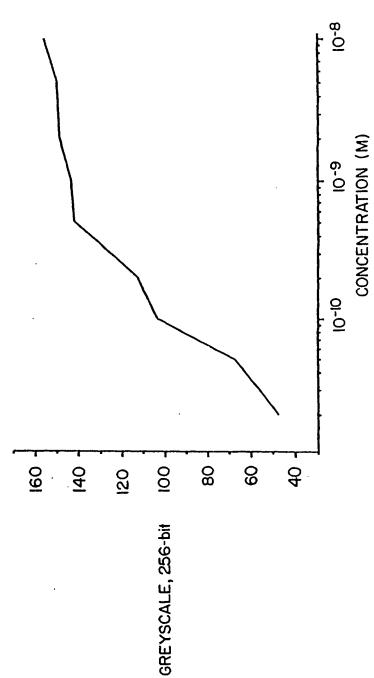
[SEQ ID NO: 57] [SEQ ID NO: 58]

N = A (complementary), G,C,T (mismatched)











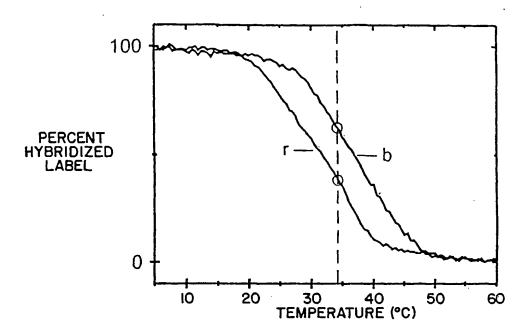


FIG.35B

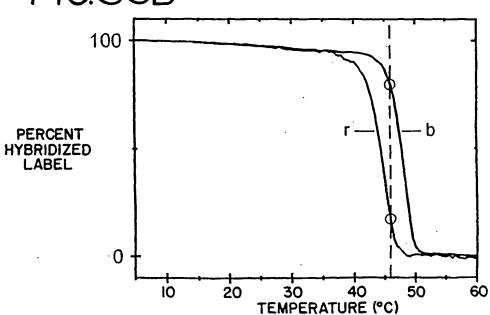


FIG. 36A

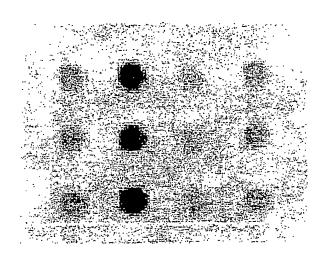
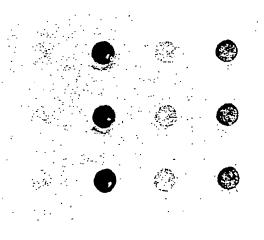


FIG. 36B



C A T G

FIG.37A

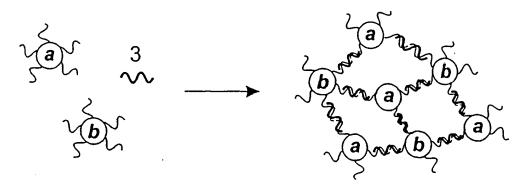
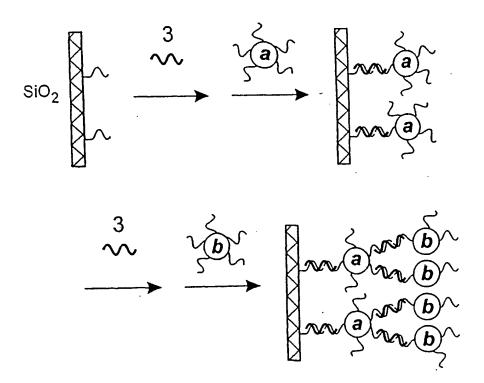
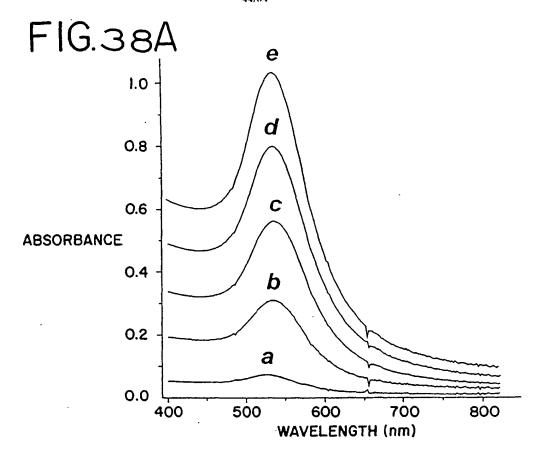


FIG.37B





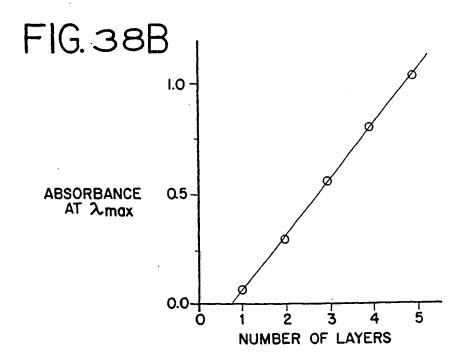


FIG. 39A

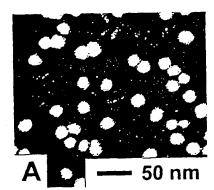
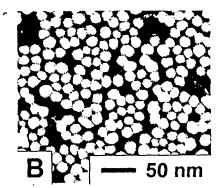
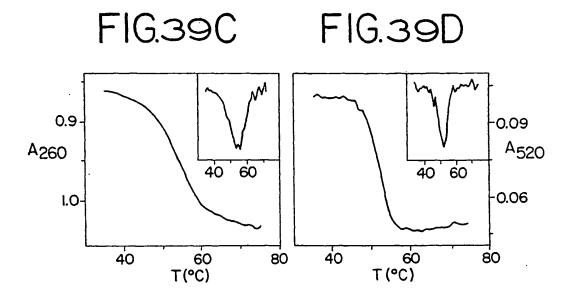
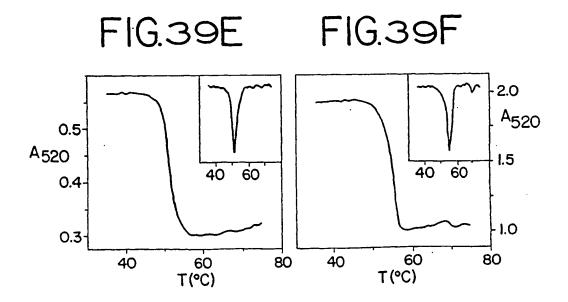
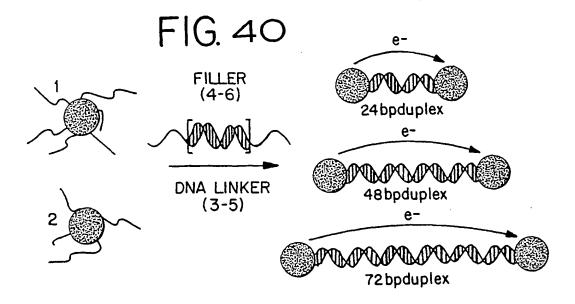


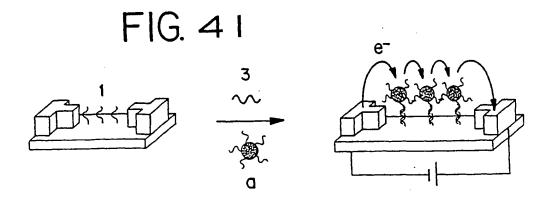
FIG.39B











 R_i

a = H

 $b = (iPr)_2NP(OCH_2CH_2CN)$ -

 $c1 = 5'p(A_{20})$ -TATCGTTCCATCAGCT [SEQ ID NO: 65]

 $c2 = 5'-p(A_{20})$ -TTGATCTTCCGTTCT [SEQ ID NO: 66]

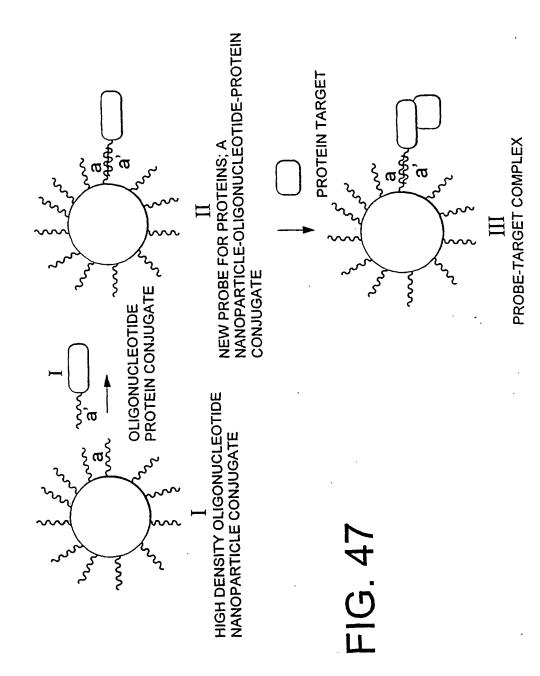
Target I = 79-mer oligonucleotide with target region:

3'-.....ATAGCAAGGTAGTCGAGCAACTAGAAAGGCAAGA.......5'
[SEQ ID NO: 67]

$$R_4$$
-(CH_2)_n R_3 R_4 -NH(CH_2)_n R_3

 R_3 = hydrogen, an alkyl group, an aryl group, or a substituted alkyl or aryl group

 R_4 = an attached oligonucleotide or modified oligonucleotide



GLASS PLATE WITH THREE DIFFERENT PROTEINS IMMOBLIZED ON THE SURFACE, ONE OF WHICH BINDS TO THE PROTEIN IN PROBE II

3

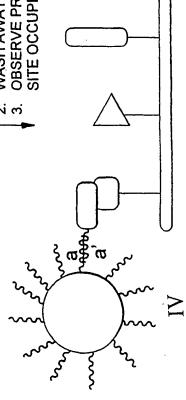
E OF WHICH

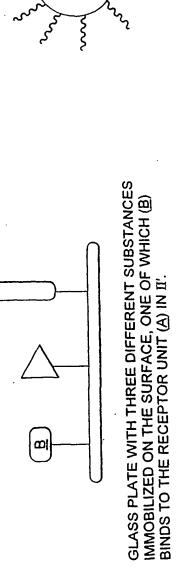
II

NEW PROBE FOR PROTEINS; A

NANOPARTICLE-OLIGONUCLEOTIDE-PROTEIN
CONJUGATE







۶ٍ **ه**ڴ۪ٟ **ٔ ٔه**



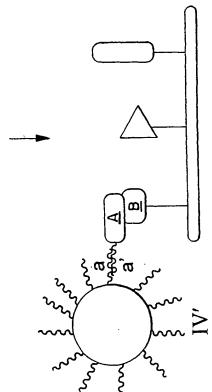
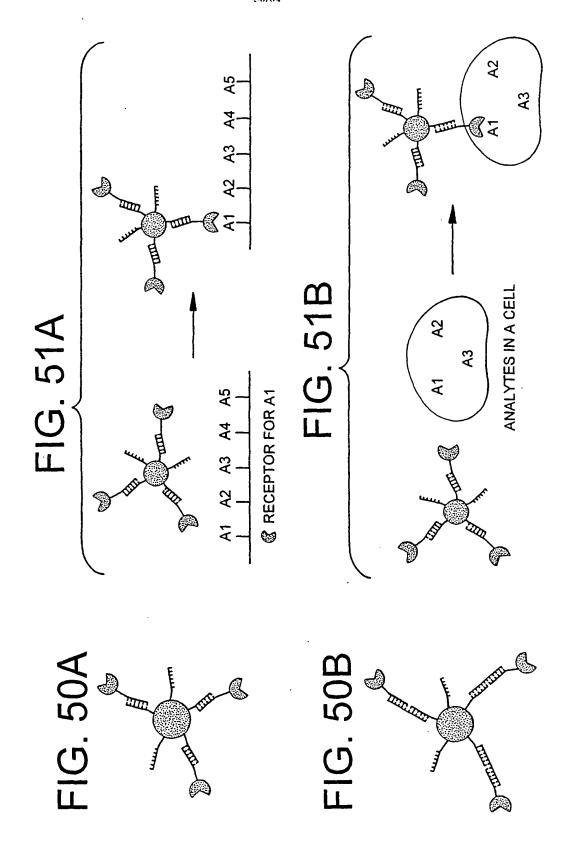
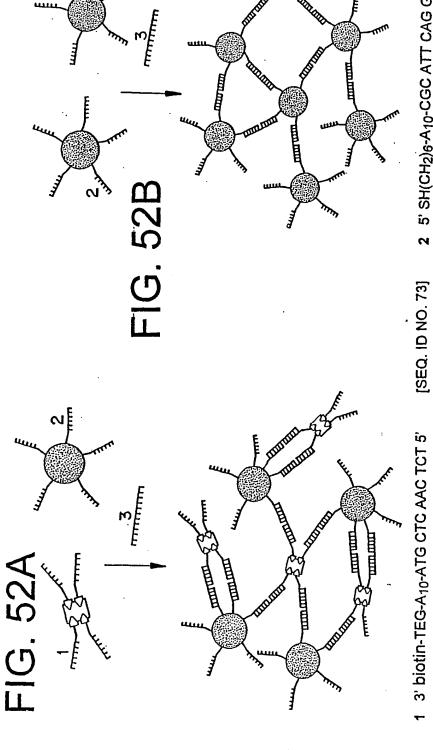


FIG. 49





2 5' SH(CH₂)6-A₁₀-CGC ATT CAG GAT 3' 3 5' TAC GAG TTG AGA ATC CTG AAT GCG 3'

[SEQ. ID NO. 74]

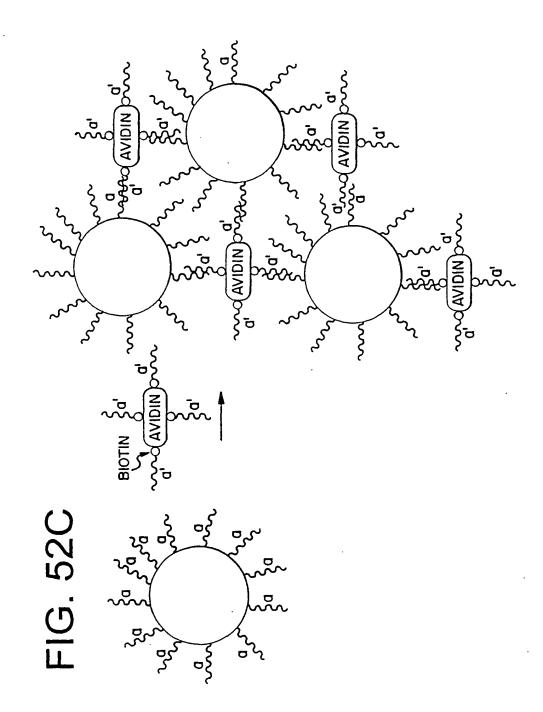
4 3' SH(CH₂)₃-A₁₀-ATG CTC AAC TCT 5'

(3) 13 nm Au NANOPARTICLES

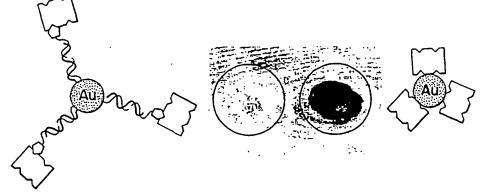
🎒 13 nm Au NANOPARTICLES 🛚 💢 STREPTAVIDIN

5' TAC GAG TTG AGA ATC CTG' AAT GCG 3' [SEQ. ID NO. 75]

5' SH(CH₂)6-A₁₀-CGC ATT CAG GAT 3'

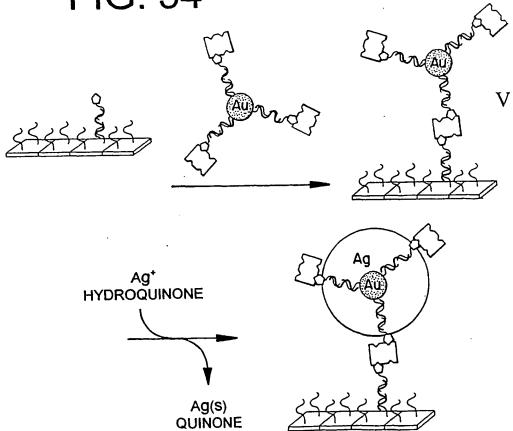


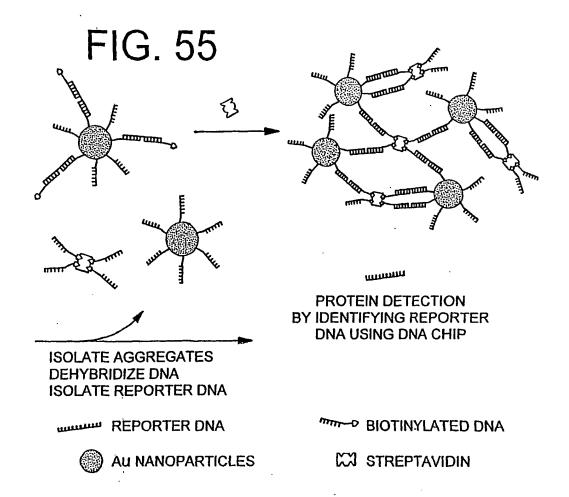


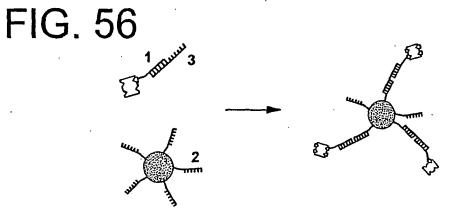


AU COLLOID / DNA / STREPTAVIDIN VS. AU COLLOID / STREPTAVIDIN CONJUGATE CONJUGATE

FIG. 54







1 3' BIOTIN-TEG-A₁₀-ATG CTC AAC TCT 5'

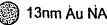
[SEQ.ID NO: 73]

2 5' SH(CH₂)₆-A₁₀-CGC ATT CAG GAT 3'

[SEQ. ID NO: 74]

3 5' TAC GAG TTG AGA ATC CTG AAT GCG 3'

[SEQ. ID NO: 75]



13nm Au NANOPARTICLES



LINKER DNA

™ BIOTINYLATED DNA

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FIG. 57A

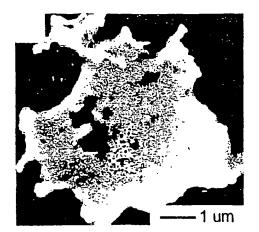


FIG. 57B

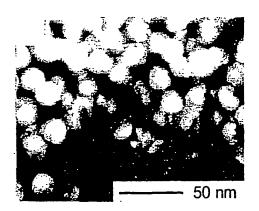


FIG. 58A

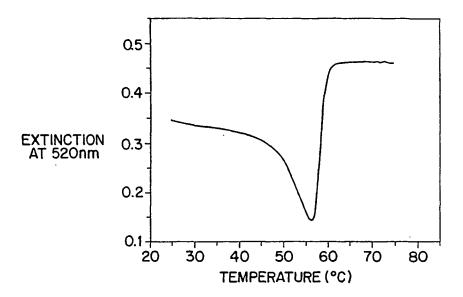


FIG. 58B

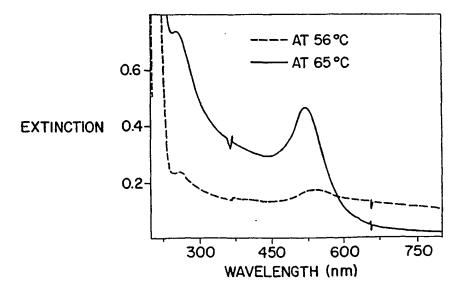


FIG. 59

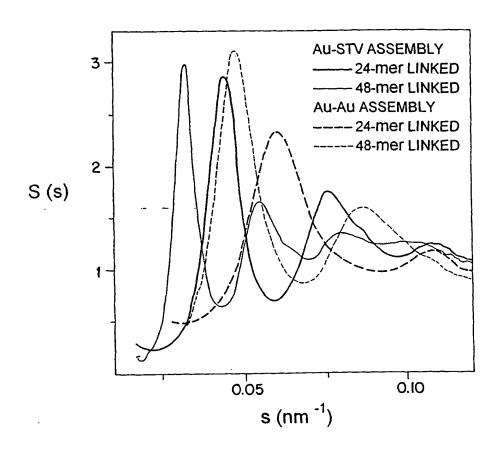
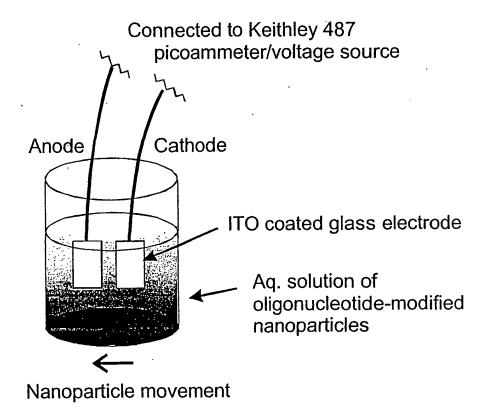


Figure 60

Scheme 1



SEQUENCE LISTING

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 Letsinger, Robert L.
 Mucic, Robert C.
 Storhoff, James J.
 Elghanian, Robert
 Taton, Thomas A.

<120> MANOPARTICLES HAVING OLIGONUCLEOTIDES ATTACHED THERETO AND USES THEREFOR

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<140> Not Yet Assigned

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<150> PCT/US97/12783

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<150> 09/240,755

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cctatcgacc atgct

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211>		
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:220>		
	Description of bubilisis Lamenas words	
2237	Description of Artificial Sequence:random synthetic sequence	
	plwengere heddeuce	

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17

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